

50

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/524,426
Source: PU/10
Date Processed by STIC: 1/30/06

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 01/23/2006

PATENT APPLICATION: US/10/524,426

TIME: 09:14:26

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\01232006\J524426.raw

3 <110> APPLICANT: Li, Limin
 4 Aghdasi, Bahman
 6 <120> TITLE OF INVENTION: MAMMALIAN GENES INVOLVED IN RAPAMYCIN RESISTANCE
 AND
 7 TUMORGENESIS: RAPR7 GENES
 9 <130> FILE REFERENCE: 70017.11USWO
 11 <140> CURRENT APPLICATION NUMBER: US 10/524,426
 12 <141> CURRENT FILING DATE: 2005-02-15
 14 <150> PRIOR APPLICATION NUMBER: PCT/US2003/026073
 15 <151> PRIOR FILING DATE: 2003-08-15
 17 <150> PRIOR APPLICATION NUMBER: US 60/404,311
 18 <151> PRIOR FILING DATE: 2002-08-15
 20 <160> NUMBER OF SEQ ID NOS: 23
 22 <170> SOFTWARE: PatentIn version 3.3
 24 <210> SEQ ID NO: 1
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 27 <213> ORGANISM: Murine
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 58 <223> OTHER INFORMATION: n is a, c, g, or t
 60 <220> FEATURE:
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63 <223> OTHER INFORMATION: n is a, c, g, or t
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66 <221> NAME/KEY: misc_feature
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76 <221> NAME/KEY: misc_feature
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81 <221> NAME/KEY: misc_feature
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110 <220> FEATURE:
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      155 tgcccttttg caggttctct tactgaccat cccacactgc cccacacatc ctcccctatg      180
W--> 157 caccccaact ntgagccctt cctgtctcagt aagtctgtag acttggtggg tatattggnc      240
      159 tcattgagac tgcaggccct tggagggcag gctctgacct gcagtaagat gtgtgagtga      300
W--> 161 tactcagcac acantaggtg gataaatacc cccacagtag gtgggtagtg agccctgtga      360
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      167 ctcccggtt cctcctgga cctgtgtgac tctgaggaac ttggggaatt cctaacctcc      540
      169 cctttcaact gagcccttgg ctcttggagt tagccacaac ctaactactc aggtccctcc      600
      171 aacaagggga ctgtgtctgt ggctggatga ctcatgcaca ctgctccatc ccgcaatctt      660
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      179 acagcaagtt caaagttcct gaggtgggaa tgcgcttgac acaacggaga cctgagaaga      900
W--> 181 acacagcaaa ggccgtgtta catttgtctg ngactccagc ccccaaggat ctggtcagga      960
W--> 183 cagacatngc gaggactcac ctggataatc cagagccatg gcccatacna ngnntncttc      1020
W--> 185 tttttttttt ttcttctttt tttctttttt tctttttttg nnnnnngccc caagacaggc      1080
W--> 187 tttctttgng tagccccggc tgttttgga ctnactntgt agaccaaact ggcctgngaa      1140
W--> 189 ctcacagaga tcctcctgnc tttgncctnc gagtacaagg gttaaaagcc tgagccanta      1200
W--> 191 ccactggcca ggctaactaa ggttcttaac tttttaagna ttatttttct ttcttatgta      1260
      193 tgtgtatatg ggggagggga tgcacaaggg catggggggg ggggtccctgc agaagtccga      1320
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      199 ctctaggccc aatgtctggt tttgttttgt tttgttttgt tttgttttgt tttgttttgt      1500
      201 tttgtatttg ggggtttttg tttgtctgtt tgggtgggtt gtttgggttt tcttgagaca      1560
      203 gggtttctct gtatagccct ggctgtcctg gaactcactc tatagactag gctggcctcg      1620

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205 aactcagaaa tctctctgcc tctgctctcc aagtgtctgag attaaaggcc cgtgccacca 1680
 207 ctgccccgacg ccaatgtctg tattttattc atctctgcag aatctctttt gtctcctaac 1740
 209 ggaacatcat cccagattct gggaagtaca ctgaagacaa tgggggtgggt gttgtttctc 1800
 W--> 211 **tcctatgccc ttacatnct ccctacctat ttcagatgta accatgatct accagctcat** 1860
 213 cacaggccac agcttaaacc tccctc 1886
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 217 <211> LENGTH: 2856
 218 <212> TYPE: DNA
 219 <213> ORGANISM: Murine
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 226 cacatctcca ccatcaacca cacccttcca tctttctctt catctgacac atatcttcca 180
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 234 tccgagatgg aagagaagag gcgaaaatat tccatcagca gcgacaactc tgataccact 420
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 238 tccggctggc cccggcagaa cgagaagaag ccctcagagg ttttcgggac agacttgatc 540
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 242 gacccgtggc gacaagaatg ggagaaagg gtgcaggtac ctgctggagc ggaggccatt 660
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 246 gatagcccca cacttggtga gggtgcccat cctgactggc caggaggcag ccgctacgac 780
 248 ctggatgaga tcgatgcgta ctggttgga cttctcaact cggagctcaa ggagatggag 840
 250 aagcccagac tggatgagct aacgttagag cgtgttctag aggagctaga gacattgtgc 900
 252 caccagaata tggcacaggc cattgagaca caggaggggc tgggcatcga gtacgacgag 960
 254 gacgttgtct gcgacgtgtg ccgttcccct gaaggcgagg atggcaacga gatggtcttc 1020
 256 tgtgacaaat gcaatgtctg tgtgcaccag gcatgctacg ggatcctcaa ggtgcctacg 1080
 258 ggcagctggc tgtgccggac ctgtgccttg ggagtcacgc ctaagtgcct gctctgcccc 1140
 260 aagcgaggag gagccctgaa gccactaga agtgggacca agtgggtaca cgtcagctgt 1200
 262 gccctgtgga ttcctgaggt cagcattggc tgtccagaga agatggagcc cattaccaag 1260
 264 atctcgcata ttccggccag ccgctgggac ctgtcctgca gcctctgcaa ggagtgaca 1320
 266 ggtacctgca tccagtgttc catgccttcc tgcacacag cattccacgt tacgtgcgcc 1380
 268 tttgaccgag gcctggaaat gcggactata ttagctgaca atgacgaggt caagttcaag 1440
 270 tcactttgcc aggagcacag tgacgggggc cctcggagtg agcctacttc tgagcctgtg 1500
 272 gagcccagcc aggcctgtga ggatctggaa aagggtacct tacgcaagca gcggctgcag 1560
 274 cagctggaag aaaacttcta tgagctagtg gagccagctg aggtggctga acggctagac 1620
 276 ctggctgagg cactggtgga cttcatctac cagtactgga agttgaagcg gagagctaata 1680
 278 gccaaaccagc cgctgttgac gcccaagact gacgaggtgg acaacctggc ccaacaggaa 1740
 280 caggatgtcc tctatcgacg cctgaagctt ttcaccacc tgcggcagga cctggagagg 1800
 282 gtaaggaacc tgtgctacat ggtgacaaga cgggagagaa cgaaacacac catctgtaaa 1860
 284 cttcaggagc agatattcca tctacagatg aaacttattg agcaagacct ttgcagagag 1920
 286 cttctgtgga ggaggtcaaa gggcaagaag aatgattcaa aaaggaaagg ccgagagggg 1980
 288 cccaagggca gcagccctga gaagaaagag aaagtgaagg ctgggcccga gtctgtgctg 2040
 290 gggcagctgg gtctatccac ctctgtcccc atcgacggca ctttcttcaa cagctggttg 2100
 292 gcacagtcgg ttcagatcac agcagaggac atggccatga gcgagtggtc tttgaacagt 2160
 294 gggcaccggg aggatcctgc tccaggtctg ctgtcagagg aattgctaca agatgaggag 2220
 296 acgctgctca gcttcatgag ggaccctcgc ctacgacctg gtgacctgc cagaaaggcc 2280
 298 cgaggccgca ctgcctgcc tgccaagaag aaacctccc cgctgcagga tgggcccagt 2340

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302 caaggaccac ccatgaggaa gccaccacgg aggacgtctt ctcatttgcc gtccagccct      2460
304 gcagctgggg actgtccagt cccagcaaca ctggaaagcc ctccaccact ggccctccgag      2520
306 atactagaca agacagcccc catggcttcc gacttaaata tccaagtgcc tggccctaca      2580
308 gtgagcccca aacccttggg caggctccgg ccaccccgag agatgaaggt cagtcggaaa      2640
310 tctccgggtg ctagatccga tgctgggaca ggactaccgt ctgctgtggc cgagaggcca      2700
312 aaggtcagcc tgcattttga caccgaggct gacggctact tctctgatga ggagatgagc      2760
314 gattctgagg tagaggcaga ggacagtggg gtacaacgag cttccaggga ggcaggggca      2820
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320 <211> LENGTH: 951
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322 <213> ORGANISM: Murine
324 <400> SEQUENCE: 3
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331                               20                               25                               30
334 Ser Leu Asn Leu Pro Pro Leu Cys His Ile Ser Thr Ile Asn His Thr
335                               35                               40                               45
338 Leu Pro Ser Phe Ser Ser Ser Asp Thr Tyr Leu Pro Thr Leu Gln Ser
339                               50                               55                               60
342 Ser Asn Lys Gln Thr Leu Lys Ala Thr Gly Pro Gly Tyr Pro Met Glu
343 65                               70                               75                               80
346 Asn Asp Gln Arg Lys Asn Thr Cys Ser Leu Val Arg Gln Glu Gly Phe
347                               85                               90                               95
350 Lys Gly Val Thr Leu His Ala Glu Ala Leu Pro Thr Glu Gly Ala Pro
351                               100                              105                              110
354 Pro Pro Pro Pro His Leu Gln Asp Ser Glu Met Glu Glu Lys Arg Arg
355                               115                              120                              125
358 Lys Tyr Ser Ile Ser Ser Asp Asn Ser Asp Thr Thr Asp Gly His Val
359                               130                              135                              140
362 Thr Ser Thr Ser Ala Ser Arg Cys Ser Lys Leu Pro Ser Ser Thr Lys
363 145                              150                              155                              160
366 Ser Gly Trp Pro Arg Gln Asn Glu Lys Lys Pro Ser Glu Val Phe Arg
367                               165                              170                              175
370 Thr Asp Leu Ile Thr Ala Met Lys Ile Pro Asp Ser Tyr Gln Leu Ser
371                               180                              185                              190
374 Pro Asp Asp Tyr Tyr Ile Leu Ala Asp Pro Trp Arg Gln Glu Trp Glu
375                               195                              200                              205
378 Lys Gly Val Gln Val Pro Ala Gly Ala Glu Ala Ile Pro Glu Pro Val
379                               210                              215                              220
382 Val Arg Leu Leu Pro Pro Leu Lys Gly Pro Pro Thr Gln Met Ser Pro
383 225                              230                              235                              240
386 Asp Ser Pro Thr Leu Gly Glu Gly Ala His Pro Asp Trp Pro Gly Gly
387                               245                              250                              255
390 Ser Arg Tyr Asp Leu Asp Glu Ile Asp Ala Tyr Trp Leu Glu Leu Leu
391                               260                              265                              270
394 Asn Ser Glu Leu Lys Glu Met Glu Lys Pro Glu Leu Asp Glu Leu Thr

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:1; N Pos. 1061,1062,1063,1064,1065,1089,1113,1117,1137,1159,1165,1168

Seq#:1; N Pos. 1198,1239,1387,1818

VERIFICATION SUMMARY

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L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:300
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:360
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:900
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:960
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1020
L:187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1080
L:189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1140
L:191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1200
L:197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1380
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1800